

EBI Dbfetch

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ID   AA689524    standard; mRNA; EST; 544 BP.
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AC   AA689524;
XX
SV   AA689524.1
XX
DT   19-DEC-1997 (Rel. 54, Created)
DT   03-MAR-2000 (Rel. 62, Last updated, Version 3)
XX
DE   ns66e01.r1 NCI_CGAP_Pr22 Homo sapiens cDNA clone IMAGE:1188600 5' similar
DE   to SW:CNI_DROME P49858 CORNICHON PROTEIN. ;, mRNA sequence.
XX
KW   EST.
XX
OS   Homo sapiens (human)
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
OC   Eutheria; Primates; Catarrhini; Hominidae; Homo.
XX
RN   [1]
RP   1-544
RA   NCI-CGAP;
RT   "National Cancer Institute, Cancer Genome Anatomy Project (CGAP), Tumor
RT   Gene Index http://www.ncbi.nlm.nih.gov/ncicgap";
RL   Unpublished.
XX
DR   RZPD; IMAGp998K012985.
DR   UNILIB; 34; 910.
XX
CC   Insert Length: 484    Std Error: 0.00
CC   Seq primer: -28m13 rev1 ET from Amersham
CC   High quality sequence stop: 499.
XX
CC   On Nov 6, 1997 this sequence version replaced gi:947554.
CC   Contact: Robert Strausberg, Ph.D.
CC   Tel: (301) 496-1550
CC   Email: Robert\_Strausberg@nih.gov
CC   Tissue Procurement: Michael J. Brownstein, M.D., Ph.D., Michael R.
CC   Emmert-Buck, M.D., Ph.D.
CC   cDNA Library Preparation: M. Bento Soares, Ph.D.
CC   cDNA Library Arrayed by: Greg Lennon, Ph.D.
CC   DNA Sequencing by: Washington University Genome Sequencing Center
CC   Clone distribution: NCI-CGAP clone distribution information can be
CC   found through the I.M.A.G.E. Consortium/LLNL at:
CC   www-bio.llnl.gov/bbrp/image/image.html
XX
FH   Key          Location/Qualifiers
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FT                  /db_xref="RZPD:IMAGp998K012985"
FT                  /db_xref="UNILIB:34"

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FT      /mol_type="mRNA"
FT      /note="Organ: prostate; Vector: pT7T3D-Pac (Pharmacia) with
FT      a modified polylinker; 1st strand cDNA was prepared from
FT      normal prostate bulk tissue, and was then primed with a Not
FT      I - oligo(dT) primer. Double-stranded cDNA was ligated to
FT      Eco RI adaptors (Pharmacia), digested with Not I and cloned
FT      into the Not I and Eco RI sites of the modified pT7T3
FT      vector. Library is normalized, and was constructed by
FT      Bento Soares and M. Fatima Bonaldo. "
FT      /sex="male"
FT      /organism="Homo sapiens"
FT      /clone="IMAGE:1188600"
FT      /clone_lib="NCI_CGAP_Pr22"
FT      /tissue_type="normal prostate"
FT      /lab_host="DH10B"
XX
SQ

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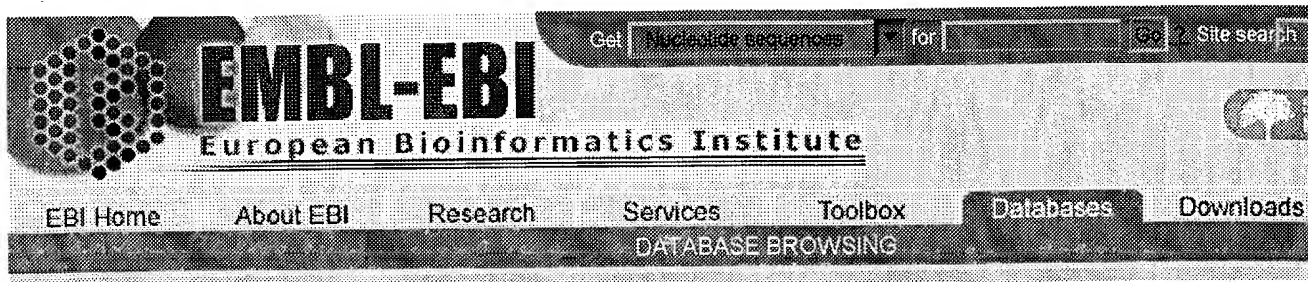
Sequence 544 BP; 165 A; 91 C; 103 G; 185 T; 0 other;
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attgtcagaa ggaaggatgg tgcaaattag ctttttatct tctagcattt ttttactacc      120
tatatggcat gatctatggt ttggtgagct cttagaacaa cacacagaag aattgggtcca      180
gttaagtgca tgcaaaaagc caccaaatga agggattcta tccagcaaga tctgtgccaa      240
gagtagcctg tggaaatctga tcagttactt taaaaaatga ctcttatctt tttaaatggt      300
tccacatttt tgcttgtgga aagactgttt tcatatgtta tactcagata aagattttta      360
atggtattac gtataaatta atataaatg attacctctg gtgttgacag gtttgaactt      420
gcacttcctt aggaacagcc ataatcctct gaatgatgca ttaattactg actgtcctag      480
tacattggaa gctttttggt ataggaactt gtagggctca ttttggtttc attgaaacag      540
tattc

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EBI Dbfetch

ID AA902726 standard; mRNA; EST; 500 BP.
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 AC AA902726;
 XX
 SV AA902726.1
 XX
 DT 09-APR-1998 (Rel. 55, Created)
 DT 03-MAR-2000 (Rel. 62, Last updated, Version 3)
 XX
 DE ok71h07.s1 NCI_CGAP_GC4 Homo sapiens cDNA clone IMAGE:1519453 3', mRNA
 DE sequence.
 XX
 KW EST.
 XX
 OS Homo sapiens (human)
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
 OC Eutheria; Primates; Catarrhini; Hominidae; Homo.
 XX
 RN [1]
 RP 1-500
 RA NCI-CGAP;
 RT "National Cancer Institute, Cancer Genome Anatomy Project (CGAP), Tumor
 RT Gene Index <http://www.ncbi.nlm.nih.gov/ncicgap>;
 RL Unpublished.
 XX
 DR RZPD; [IMAGp998D143847](#).
 DR UNILIB; [46](#); 1050.
 XX
 CC Insert Length: 1413 Std Error: 0.00
 CC Seq primer: -40m13 fwd. ET from Amersham
 CC High quality sequence stop: 375.
 XX
 CC On Jan 9, 1998 this sequence version replaced gi:936894.
 CC Contact: Robert Strausberg, Ph.D.
 CC Tel: (301) 496-1550
 CC Email: Robert.Strausberg@nih.gov
 CC Tissue Procurement: Christopher A. Moskaluk, M.D., Ph.D., Michael
 CC Emmert-Buck, M.D., Ph.D.
 CC cDNA Library Preparation: M. Bento Soares, Ph.D.
 CC cDNA Library Arrayed by: Greg Lennon, Ph.D.
 CC DNA Sequencing by: Washington University Genome Sequencing Center
 CC Clone distribution: NCI-CGAP clone distribution information can be
 CC found through the I.M.A.G.E. Consortium/LLNL at:
 CC www-bio.llnl.gov/bbrp/image/image.html
 XX
 FH Key Location/Qualifiers
 FH
 FT source 1..500
 FT /db_xref="taxon:9606"
 FT /db_xref="RZPD:IMAGp998D143847"
 FT /db_xref="UNILIB:46"

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FT      /mol_type="mRNA"
FT      /note="Vector: pT7T3D-Pac (Pharmacia) with a modified
FT      polylinker; 1st strand cDNA was prepared from 3 pooled germ
FT      cell tumors, and was then primed with a Not I - oligo(dT)
FT      primer. Double-stranded cDNA was ligated to Eco RI adaptors
FT      (Pharmacia), digested with Not I and cloned into the Not I
FT      and Eco RI sites of the modified pT7T3 vector. Library is
FT      normalized. Library was constructed by Bento Soares and M.
FT      Fatima Bonaldo. "
FT      /organism="Homo sapiens"
FT      /clone="IMAGE:1519453"
FT      /clone_lib="NCI_CGAP_GC4"
FT      /tissue_type="pooled germ cell tumors"
FT      /lab_host="DH10B"
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SQ
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Sequence 500 BP; 180 A; 81 C; 66 G; 173 T; 0 other;
taatttcaga gttttattgt attgcactaa aggaacagca ggatgggttat acaattttct      60
ctcattcagt ttgaaaatc tgtagtacct gcaaattctt aagaatacct ttaccaccag      120
attagaacag taagcataat aaccaatttc ttaataagta atgtcttaca aataaaaaaca      180
catttaaaat agctttaaat gcattcttca caagtaattc agcatatatt tttatatcat      240
gtttacttat gcttaagaat taaagcaagt atatttatta ctctgatgga aatatgggaa      300
atctctcatt catgcaatat acagggataa tattcaagcg aagggaatat tcccgtttt      360
tatttttgta aatgtatcca tatataatca tcgacatgac agatgaggaa acccatgaag      420
tttcccacta gtcagatata cattttcact tcatcagaag cacctgatat ctacagctaa      480
tttataatta gatactgttt                                         500
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